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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/668,558BDATE: 09/13/2001  
TIME: 10:26:08Input Set : D:\Seqlist.txt  
Output Set: N:\CRF3\09132001\I668558B.raw

5 <110> APPLICANT: Yen, Frances  
6 Erickson, Mary Ruth  
7 Fruebis, Joachim  
8 Bihain, Bernard  
10 <120> TITLE OF INVENTION: Methods Of Screening For Compounds That Modulate the  
11 LSR-Leptin Interaction and Their Use in the Prevention  
12 and Treatment of Obesity-Related Diseases  
14 <130> FILE REFERENCE: 70.US2.REG  
16 <140> CURRENT APPLICATION NUMBER: 09/668,558B  
17 <141> CURRENT FILING DATE: 2000-09-22  
19 <150> PRIOR APPLICATION NUMBER: 60/155,506  
20 <151> PRIOR FILING DATE: 1999-09-22  
22 <160> NUMBER OF SEQ ID NOS: 106  
24 <170> SOFTWARE: Patent.pm  
26 <210> SEQ ID NO: 1  
27 <211> LENGTH: 23187  
28 <212> TYPE: DNA  
29 <213> ORGANISM: Homo sapiens  
31 <220> FEATURE:  
32 <221> NAME/KEY: exon  
33 <222> LOCATION: 2001..2356  
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38 <222> LOCATION: 3540..3884  
39 <223> OTHER INFORMATION: exon2  
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43 <222> LOCATION: 12163..12282  
44 <223> OTHER INFORMATION: exon3  
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53 <222> LOCATION: 15765..15911  
54 <223> OTHER INFORMATION: exon5  
56 <220> FEATURE:  
57 <221> NAME/KEY: exon  
58 <222> LOCATION: 19579..19752  
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61 <220> FEATURE:  
62 <221> NAME/KEY: exon  
63 <222> LOCATION: 19899..19958  
64 <223> OTHER INFORMATION: exon7  
66 <220> FEATURE:

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69 <223> OTHER INFORMATION: exon8
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73 <222> LOCATION: 20329..20957
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79 <223> OTHER INFORMATION: exon10
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83 <222> LOCATION: 21168..21173
84 <223> OTHER INFORMATION: AATAAA
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87 <221> NAME/KEY: misc_feature
88 <222> LOCATION: 1..2000
89 <223> OTHER INFORMATION: potential 5'regulatory region
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92 <221> NAME/KEY: misc_feature
93 <222> LOCATION: 22324..23187
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104 <223> OTHER INFORMATION: downstream amplification primer 17-2 , complement
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123 <222> LOCATION: 1616..1635
124 <223> OTHER INFORMATION: downstream amplification primer 9-19 , complement
126 <220> FEATURE:
127 <221> NAME/KEY: primer_bind

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Input Set : D:\Seqlist.txt  
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128 <222> LOCATION: 1602..1621  
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154 <223> OTHER INFORMATION: downstream amplification primer 9-1 , complement  
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158 <222> LOCATION: 2470..2489  
159 <223> OTHER INFORMATION: upstream amplification primer 9-21 , complement  
161 <220> FEATURE:  
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163 <222> LOCATION: 2062..2081  
164 <223> OTHER INFORMATION: downstream amplification primer 9-21  
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167 <221> NAME/KEY: primer\_bind  
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171 <220> FEATURE:  
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178 <222> LOCATION: 3775..3792  
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184 <223> OTHER INFORMATION: downstream amplification primer 99-4558 , complement  
186 <220> FEATURE:  
187 <221> NAME/KEY: primer\_bind  
188 <222> LOCATION: 4902..4920

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Input Set : D:\Seqlist.txt  
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189 <223> OTHER INFORMATION: upstream amplification primer 99-14419 , complement  
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197 <221> NAME/KEY: primer\_bind  
198 <222> LOCATION: 6638..6655  
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201 <220> FEATURE:  
202 <221> NAME/KEY: primer\_bind  
203 <222> LOCATION: 7072..7089  
204 <223> OTHER INFORMATION: downstream amplification primer 99-4577 , complement  
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207 <221> NAME/KEY: primer\_bind  
208 <222> LOCATION: 7995..8012  
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213 <222> LOCATION: 8576..8593  
214 <223> OTHER INFORMATION: downstream amplification primer 99-4559 , complement  
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222 <221> NAME/KEY: primer\_bind  
223 <222> LOCATION: 10023..10040  
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233 <222> LOCATION: 10546..10563  
234 <223> OTHER INFORMATION: downstream amplification primer 99-4560 , complement  
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244 <223> OTHER INFORMATION: downstream amplification primer 99-14411  
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248 <222> LOCATION: 11972..11990  
249 <223> OTHER INFORMATION: upstream amplification primer 99-4561

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254 <223> OTHER INFORMATION: downstream amplification primer 99-4561 , complement
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258 <222> LOCATION: 12005..12023
259 <223> OTHER INFORMATION: upstream amplification primer 9-4
261 <220> FEATURE:
262 <221> NAME/KEY: primer_bind
263 <222> LOCATION: 12417..12436
264 <223> OTHER INFORMATION: downstream amplification primer 9-4 , complement
266 <220> FEATURE:
267 <221> NAME/KEY: primer_bind
268 <222> LOCATION: 14102..14119
269 <223> OTHER INFORMATION: upstream amplification primer 99-4562
271 <220> FEATURE:
272 <221> NAME/KEY: primer_bind
273 <222> LOCATION: 14543..14563
274 <223> OTHER INFORMATION: downstream amplification primer 99-4562 , complement
276 <220> FEATURE:
277 <221> NAME/KEY: primer_bind
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279 <223> OTHER INFORMATION: upstream amplification primer 99-3149
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283 <222> LOCATION: 14848..14865
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287 <221> NAME/KEY: primer_bind
288 <222> LOCATION: 14748..14767
289 <223> OTHER INFORMATION: upstream amplification primer 9-22
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292 <221> NAME/KEY: primer_bind
293 <222> LOCATION: 15198..15218
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299 <223> OTHER INFORMATION: upstream amplification primer 9-24
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303 <222> LOCATION: 15333..15351
304 <223> OTHER INFORMATION: downstream amplification primer 9-24 , complement
306 <220> FEATURE:
307 <221> NAME/KEY: primer_bind
308 <222> LOCATION: 15002..15019
309 <223> OTHER INFORMATION: upstream amplification primer 9-5
311 <220> FEATURE:

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Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/668,558B

DATE: 09/13/2001

TIME: 10:26:09

Input Set : D:\Seqlist.txt

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L:1460 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:1462 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:1463 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:1587 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2  
L:1596 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2  
L:1642 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2  
L:1854 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:1863 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:2105 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6  
L:2114 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6  
L:2585 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:10  
L:2974 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:14  
L:4421 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:92  
L:4437 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:93  
L:4464 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:95  
L:4502 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:98  
L:4518 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:99  
L:4534 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:100